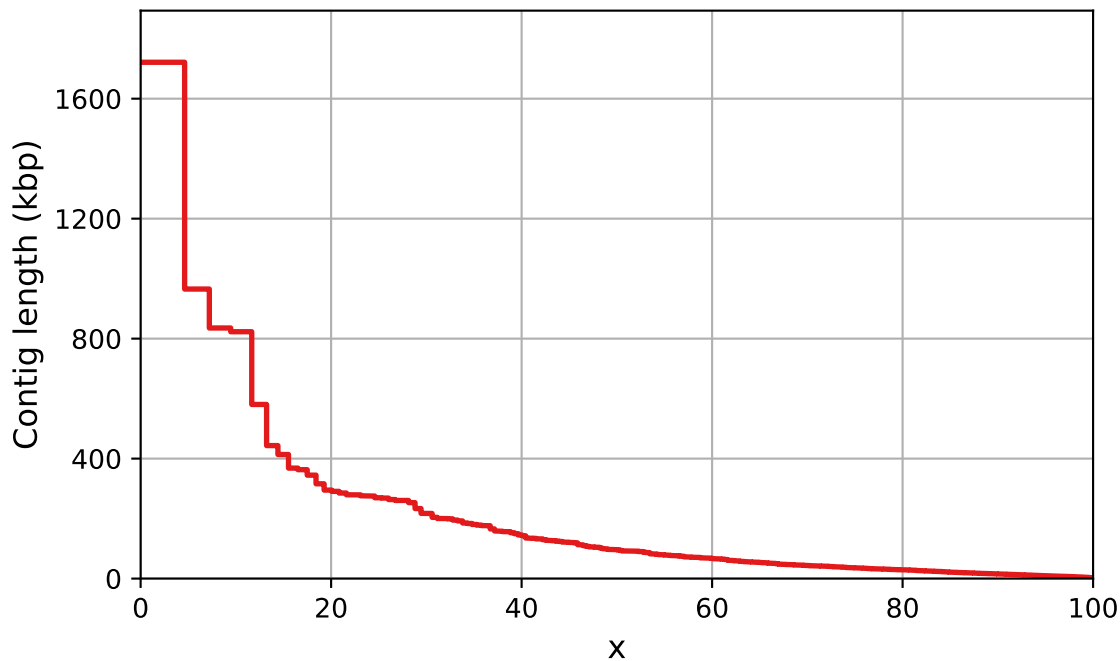


Report

	genome.nextpolish
# contigs (>= 0 bp)	944
# contigs (>= 1000 bp)	939
# contigs (>= 5000 bp)	829
# contigs (>= 10000 bp)	609
# contigs (>= 25000 bp)	343
# contigs (>= 50000 bp)	169
Total length (>= 0 bp)	37246261
Total length (>= 1000 bp)	37242534
Total length (>= 5000 bp)	36832506
Total length (>= 10000 bp)	35216957
Total length (>= 25000 bp)	30952094
Total length (>= 50000 bp)	24835972
# contigs	944
Largest contig	1721387
Total length	37246261
GC (%)	57.66
N50	96431
N90	15498
auN	252395.4
L50	79
L90	476
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	84 + 13 part

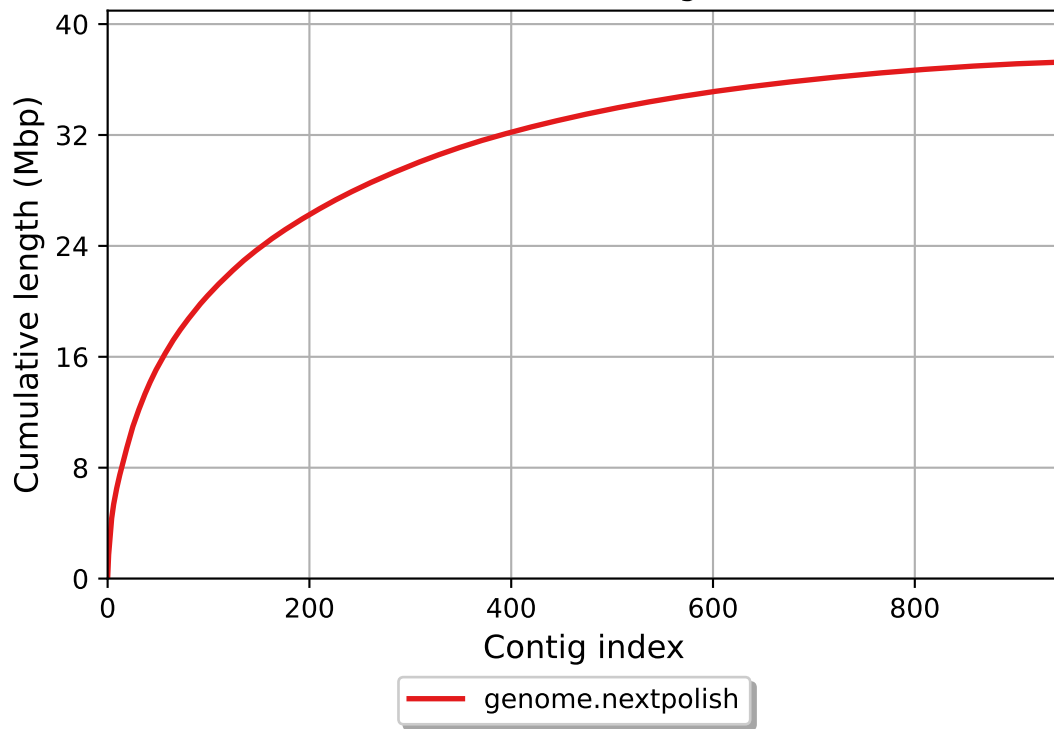
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

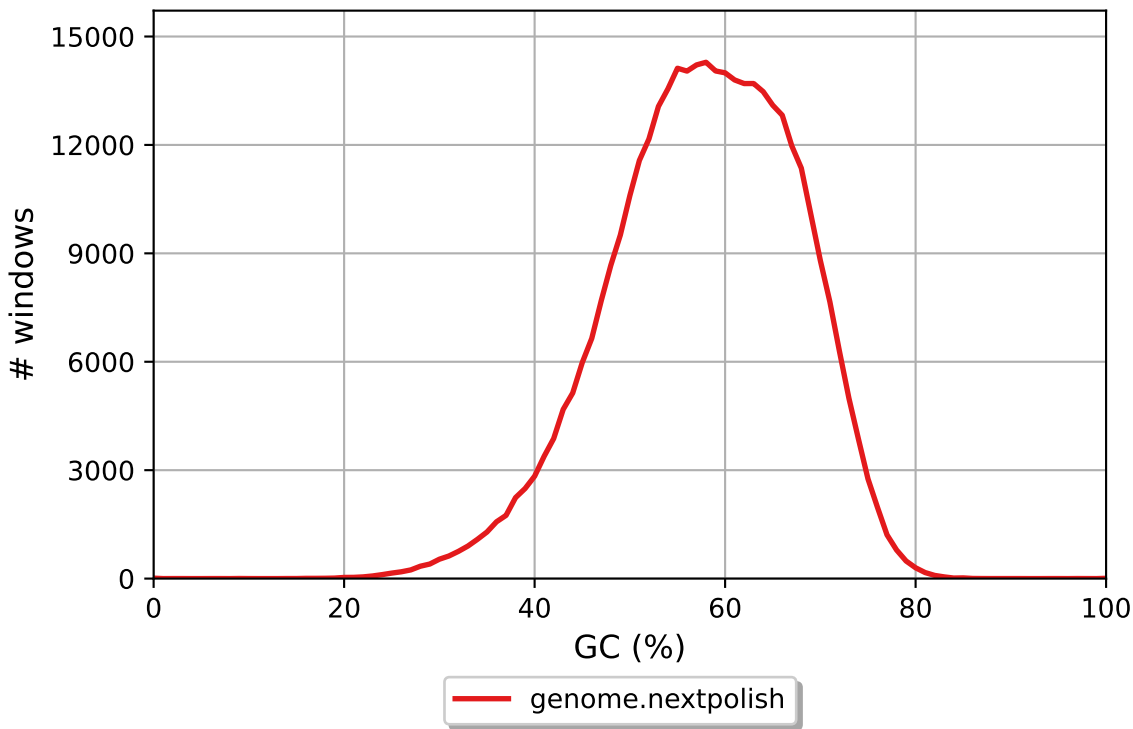


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

